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<110> Dai, Ken-Shwo	
<120> HUMAN ARL-RELATED GENE VARIANTS ASSOCIATED WITH CANCER	
<130> U 014798-3	
<140> 10/653,681 <141> 2003-09-02	
<160>. 4	
<170> PatentIn version 3.2	
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att gtg ggc ctg ggc act tgg aag tct cct ctc ggc aaa gtg aaa gaa Ile Val Gly Leu Gly Thr Trp Lys Ser Pro Leu Gly Lys Val Lys Glu 15 20 25 30	159
gca gtg aag gtg gcc att gat gca gga tat cgg cac att gac tgt gcc Ala Val Lys.Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala 35 40 45	207
Ala Val Lys. Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala	207 255
Ala Val Lys. Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala 35 40 45 tat gtc tat cag aat gaa cat gaa gtg ggg gaa gcc atc caa gag aag Tyr Val Tyr Gln Asn Glu His Glu Val Gly Glu Ala Ile Gln Glu Lys	
Ala Val Lys.Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala 35 40 45 tat gtc tat cag aat gaa cat gaa gtg ggg gaa gcc atc caa gag aag Tyr Val Tyr Gln Asn Glu His Glu Val Gly Glu Ala Ile Gln Glu Lys 50 55 60 atc caa gag aag gct gtg aag cgg gag gac ctg ttc atc gtc agc aag Ile Gln Glu Lys Arg Glu Asp Leu Phe Ile Val Ser Lys	255
Ala Val Lys Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala 35 tat gtc tat cag aat gaa cat gaa gtg ggg gaa gcc atc caa gag aag Tyr Val Tyr Gln Asn Glu His Glu Val Gly Glu Ala Ile Gln Glu Lys 50 atc caa gag aag gct gtg aag cgg gag gac ctg ttc atc gtc agc aag Ile Gln Glu Lys Arg Glu Asp Leu Phe Ile Val Ser Lys 65 ttg tgg ccc act tcc aga tcg aga agc tct tgaacaaacc tggactgaaa Leu Trp Pro Thr Ser Arg Ser Arg Ser Ser	255

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Tyr Gln Asn Glu His Glu Val Gly Glu Ala Ile Gln Glu Lys Ile Gln 50 55 60

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gtt acg gcc tac agc ccc ctg ggc tct ccg gat aga cct tgg gcc aag Val Thr Ala Tyr Ser Pro Leu Gly Ser Pro Asp Arg Pro Trp Ala Lys 210 215 220	735
cca gaa gac cct tcc ctg ctg gag gat ccc aag att aag gag att gct Pro Glu Asp Pro Ser Leu Leu Glu Asp Pro Lys Ile Lys Glu Ile Ala 225 230 235	783
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Glu Lys Ala Val Lys Arg Glu Asp Leu Phe Ile Val Ser Lys Leu Trp 65 70 75 80

Pro Thr Phe Phe Glu Arg Pro Leu Val Arg Lys Ala Phe Glu Lys Thr

85

90

95

Leu Lys Asp L'eu Lys Leu Ser Tyr Leu Asp Val Tyr Leu Ile His Trp

Pro Gln Gly Phe Lys Ser Gly Asp Asp Leu Phe Pro Lys Asp Asp Lys 120

Gly Asn Ala Ile Gly Gly Lys Ala Thr Phe Leu Asp Ala Trp Glu Ala 130 135 140

Met Glu Glu Leu Val Asp Glu Gly Leu Val Lys Ala Leu Gly Val Ser

Asn Phe Ser His Phe Gln Ile Glu Lys Leu Leu Asn Lys Pro Gly Leu

Lys Tyr Lys Pro Val Thr Asn Gln Val Glu Cys His Pro Tyr Leu Thr 185

Gln Glu Lys Leu Ile Gln Tyr Cys His Ser Lys Gly Ile Thr Val Thr

Ala Tyr Ser Pro Leu Gly Ser Pro Asp Arg Pro Trp Ala Lys Pro Glu 210 215 220

Asp Pro Ser Leu Leu Glu Asp Pro Lys Ile Lys Glu Ile Ala Ala Lys 230 235

His Ser Pro Ser Leu 245